

IFW16

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/054,665A

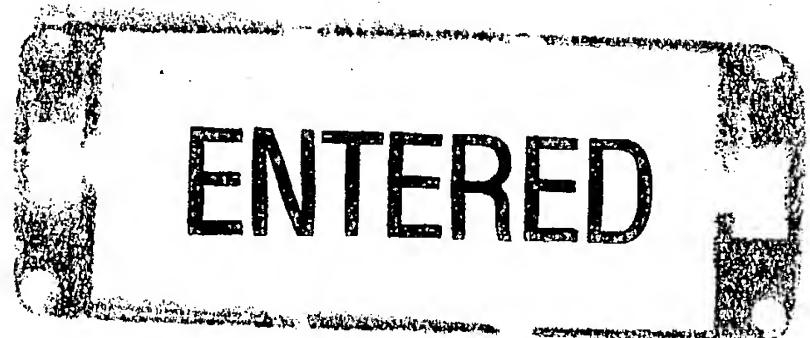
DATE: 09/24/2004

TIME: 11:46:26

Input Set : A:\875007US2.txt

Output Set: N:\CRF4\09242004\J054665A.raw

4 <110> APPLICANT: Engelhardt, John F.
5 Duan, Dongshen
6 University of Iowa Research Foundation
9 <120> TITLE OF INVENTION: Adeno-associated virus vectors
12 <130> FILE REFERENCE: 875.007US2
C--> 14 <140> CURRENT APPLICATION NUMBER: US/10/054,665A
15 <141> CURRENT FILING DATE: 2002-01-22
17 <150> PRIOR APPLICATION NUMBER: US 60/086,166
18 <151> PRIOR FILING DATE: 1998-05-20
20 <150> PRIOR APPLICATION NUMBER: US 09/276,625
21 <151> PRIOR FILING DATE: 1999-03-25
23 <160> NUMBER OF SEQ ID NOS: 14
25 <170> SOFTWARE: FastSEQ for Windows Version 4.0
27 <210> SEQ ID NO: 1
28 <211> LENGTH: 20
29 <212> TYPE: DNA
30 <213> ORGANISM: Adeno-associated virus
32 <400> SEQUENCE: 1
33 cgggggtcgt tgggcggtca
35 <210> SEQ ID NO: 2
36 <211> LENGTH: 19
37 <212> TYPE: DNA
38 <213> ORGANISM: Adeno-associated virus
40 <400> SEQUENCE: 2
41 gggcgagcc tatggaaaa
43 <210> SEQ ID NO: 3
44 <211> LENGTH: 505
45 <212> TYPE: DNA
46 <213> ORGANISM: Artificial Sequence
48 <220> FEATURE:
49 <223> OTHER INFORMATION: A synthetic consensus sequence
51 <400> SEQUENCE: 3
52 cgggggtcgt tgggcggtca gccaggcggg ccatttaccc taagttatgt a
53 ggcatgcaag ctcgaattca tcggtagata agtagcatgg cgggttaatc a
54 aggaaccctt agtgatggag ttggccactc cctctctgcg cgctcgctcg c
55 ccgggcgacc aaaggtcgcc cgacgccccg gctttgcccg ggccggctca g
56 gagcgcgcag ctgcgcgctc gtcgcctcac tgaggccgcc cgggcaaagc c
57 ggcgacctt ggtcgccccg ctcagcgag cgagcgagcg cgcagagagg g
58 ctccatcaact aggggttcct ttagttaat gattaacccg ccatgctact t
59 ttgcatgcat gtgagcaaaa ggccagcaaa aggccaggaa ccgtaaaaag g
60 tggcgaaaa ccataggctc cgccc
65 <210> SEQ ID NO: 4
66 <211> LENGTH: 272



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67 <212> TYPE: DNA
 68 <213> ORGANISM: AAV circular intermediate, clone p81
 70 <400> SEQUENCE: 4
 71 gcatgcaagc tgttagataag tagcatggcg ggttaatcat taactacaag gaaccctag 60
 72 ttagtggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcgccaa 120
 73 aggtcgcccg acgcccggc tttgccggg cggcctcagt gagcgagcga gcgcgcagag 180
 74 agggagtgcc caactccatc actagggtt cctttagtt aatgattaac ccgcattgtct 240
 75 acttatctac cgatgaattc gagcttgcac gc 272
 77 <210> SEQ ID NO: 5
 78 <211> LENGTH: 300
 79 <212> TYPE: DNA
 80 <213> ORGANISM: AAV circular intermediate, clone p79
 82 <400> SEQUENCE: 5
 83 gcatgcaagc tgttagataag tagcatggcg ggttaatcat taactacaag gaaccctag 60
 84 ttagtggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcgccgc 120
 85 tcgcgtcgctc actgaggccg ggcgacccaa ggtcgcccg gcccggctt tgcccgccg 180
 86 gcctcagtga gcgagcgcgc gcgcagagag ggagtggcca actccatcac tagggttcc 240
 87 ttgttagttaa tgattaaccc gccatgtac ttatctaccg atgaattcga gcttgcac 300
 89 <210> SEQ ID NO: 6
 90 <211> LENGTH: 272
 91 <212> TYPE: DNA
 92 <213> ORGANISM: AAV circular intermediate, clone p1202
 94 <400> SEQUENCE: 6
 95 gcatgcaagc tgttagataag tagcatggcg ggttaatcat taactacaag gaaccctag 60
 96 ttagtggagtt ggccactccc tctctgcgcg ctcgctcgct cactgaggcc gggcgaccaa 120
 97 aggtcgcccg acgcccggc tttggcgcg cggcctcagt gagcgagcga gcgcgcagag 180
 98 agggagtgcc caactccatc actagggtt cctttagtt aatgattaac ccgcattgtct 240
 99 acttatctac cgatgaattc gagcttgcac gc 272
 101 <210> SEQ ID NO: 7
 102 <211> LENGTH: 165
 103 <212> TYPE: DNA
 104 <213> ORGANISM: Unknown
 106 <220> FEATURE:
 107 <223> OTHER INFORMATION: SEQ ID NO:1 of U.S. Patent No. 5,478,745
 109 <400> SEQUENCE: 7
 110 aggaacccct agtgatggag ttggccactc cctctctgcg cgctcgctcg ctcactgagg 60
 111 cccggcgacc aaaggtcgcc cgacgcccgg gctttggccg ggcggccta gtgagcgcg 120
 112 gagcgccgag agagggagtg gccaactcca tcactagggg ttccct 165
 114 <210> SEQ ID NO: 8
 115 <211> LENGTH: 282
 116 <212> TYPE: DNA
 117 <213> ORGANISM: rAAV circular intermediate, clone p79
 119 <400> SEQUENCE: 8
 120 ggcggccat ttaccgtaa ttatgtggcg actgcaggca tgcaagctcg aattcatcg 60
 121 tagataagta gcatggcggt ttaatcattg cctacaaaga gccccttagtg atggagtg 120
 122 ccactccctc tcttcgcgcg gcgcgcagag agggagtgcc caactccctc actagggtt 180
 123 cctggcagtt aatgattaac ccgcattgtct acttatctac agcttgcac catgtgagca 240
 124 aaaggccagc aaaaggccag gaaccgtaaa aaggccgcgt tg 282
 127 <210> SEQ ID NO: 9

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128 <211> LENGTH: 345
 129 <212> TYPE: DNA
 130 <213> ORGANISM: rAAV circular intermediate, clone p80
 132 <400> SEQUENCE: 9
 133 ggccattttac cgtaagttat gtaacgactg caggcatgca agctcgaatt catcggtaga 60
 134 taagtagcat ggccgggttaa tcattaacta caaggaaccc ctatgtatgg agttggccac 120
 135 tccctctctg cgcgctcgct cgctcgctca ggccggggcga ccaaaggctcg cccgacgccc 180
 136 gcccggcctc agcgagcgag cgagcgcgca gagagggagt ggccaactcc atcacttaggg 240
 137 gttccttgcgttaatgatt aaccggccat gctacttatac tacagcttgc atgcatgtga 300
 138 gcaaaaaggcc agcaaaaaggc caggaaccgt aaaaaggccg cgttg 345
 140 <210> SEQ ID NO: 10
 141 <211> LENGTH: 276
 142 <212> TYPE: DNA
 143 <213> ORGANISM: rAAV circular intermediate, clone p81
 145 <400> SEQUENCE: 10
 146 ggccattttac cgtaagttat gtggcgactg caggcatgca agctcgaatt catcggtaga 60
 147 taagtagcat ggccgggttaa tcattgccta caaagagccc ctatgtatgg agcccgccct 120
 148 caccgagcga gcgagcgccg agagagggag tgcccaactc catcaactagg gttccttgt 180
 149 agttaatgat taacccgcca tgctacttatac tacagcttgc atgcatgtga agcaaaaaggc 240
 150 cagcaaaaagg ccaggaaccg taaaaaggcc gcggtg 276
 152 <210> SEQ ID NO: 11
 153 <211> LENGTH: 316
 154 <212> TYPE: DNA
 155 <213> ORGANISM: rAAV circular intermediate, clone p86
 157 <400> SEQUENCE: 11
 158 ggccattttac cgtaagttat gtaacgactg caggcatgca agctcgaatt catcggtaga 60
 159 taagtagcat ggccgggttaa tcattaacta caaggaaccc ctatgtatgg agttggccac 120
 160 tccctctctg cgcgctcgct cgctcgctga ggccggcccg gcctcagcga gcgagcgagc 180
 161 ggcgcagagag ggactggcca actccatcac taggggttcc ttgttagttaa tgattaaccc 240
 162 gccatgtac ttatctacag cttgcatacg tgtgagcaaa aggccagcaa aaggccagga 300
 163 accgtaaaaa ggccgc 316
 165 <210> SEQ ID NO: 12
 166 <211> LENGTH: 208
 167 <212> TYPE: DNA
 168 <213> ORGANISM: rAAV circular intermediate, clone p87
 170 <400> SEQUENCE: 12
 171 ggccattttac cgtaagttat gtaacgactg caggcatgca agctcgaatt catcggtaga 60
 172 taagtagcat ggccgggttac tcattgccta caaagagccc ctatgtatgg aattggaaatg 120
 173 attcacccttc catgtactt atctacagct tgcatgcata tgagcaaaag gccagcaaaa 180
 174 ggccaggaac cgtaaaaagg ccgcgttg 208
 176 <210> SEQ ID NO: 13
 177 <211> LENGTH: 310
 178 <212> TYPE: DNA
 179 <213> ORGANISM: rAAV circular intermediate, clone p88
 181 <400> SEQUENCE: 13
 182 gccattttacc gtaagttatg taacgactgc aggcatgcaa gctcgaattc atcggtatgt 60
 183 aagtagcatg ggccgggttaat cattgcctac aaagagcccc tagtgtatgg gttggccact 120
 184 ccctctctgc gcgctcgctc gctggggcccg gcctcagcga gcgagcgagc ggcgcagagag 180
 185 ggagtggcca actccatcac taggggttcc ttgttagttaa tgattaaccc gccatgtac 240

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186 ttatctacag cttgcatgca tgtgagcaa aggccagcaa aaggccagga accgtaaaaa 300
187 ggccgcgttg 310
189 <210> SEQ ID NO: 14
190 <211> LENGTH: 334
191 <212> TYPE: DNA
192 <213> ORGANISM: Artificial Sequence
194 <220> FEATURE:
195 <223> OTHER INFORMATION: A synthetic portion of the consensus sequence
197 <400> SEQUENCE: 14
198 gtagataagt agcatggcggtttaatcatt aactacaagg aacccttagt gatggagttg 60
199 gccactccct ctctgcgcgc tcgctcgctc gctgaggccg ggcgaccaaa ggtcgcccga 120
200 cgcccgggct ttgcccgggc gcgcctcagtg agcgagcgag cgcgcagctg cgcgctcgct 180
201 cgctcactga ggccgccccgg gcaaagcccg ggcgtcgggc gaccttttgtt cgcccggcct 240
202 cagcgagcga gcgagcgcgc agagagggag tggccaactc catcactagg gtttccttgt 300
203 agttaatgat taacccgcca tgctacttat ctac 334

VERIFICATION SUMMARY

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L:14 M:270 C: Current Application Number differs, Replaced Current Application Number